\_\_\_\_\_\_

Sequence Listing was accepted.

See attached Validation Report.

If you need help call the Patent Electronic Business Center at (866)

217-9197 (toll free).

Reviewer: Keisha Douglas

Timestamp: [year=2008; month=1; day=17; hr=17; min=7; sec=47; ms=220; ]

\_\_\_\_\_\_

## Validated By CRFValidator v 1.0.3

Application No: 10532480 Version No: 2.0

Input Set:

Output Set:

**Started:** 2008-01-04 18:09:22.369

Finished: 2008-01-04 18:09:27.470

**Elapsed:** 0 hr(s) 0 min(s) 5 sec(s) 101 ms

Total Warnings: 69

Total Errors: 72

No. of SeqIDs Defined: 75

Actual SeqID Count: 75

Error code		Error Description	
W	213	Artificial or Unknown found in <213> in SEQ ID (	L)
E	257	Invalid sequence data feature in <221> in SEQ ID	(1)
Ε	257	Invalid sequence data feature in <221> in SEQ ID	(1)
W	213	Artificial or Unknown found in <213> in SEQ ID (2	2)
Ε	257	Invalid sequence data feature in <221> in SEQ ID	(2)
Ε	257	Invalid sequence data feature in <221> in SEQ ID	(2)
W	213	Artificial or Unknown found in <213> in SEQ ID (3	3)
Ε	257	Invalid sequence data feature in <221> in SEQ ID	(3)
E	257	Invalid sequence data feature in <221> in SEQ ID	(3)
W	213	Artificial or Unknown found in <213> in SEQ ID (	1)
E	257	Invalid sequence data feature in <221> in SEQ ID	(4)
E	257	Invalid sequence data feature in <221> in SEQ ID	(4)
W	213	Artificial or Unknown found in <213> in SEQ ID (	5)
E	257	Invalid sequence data feature in <221> in SEQ ID	(5)
E	257	Invalid sequence data feature in <221> in SEQ ID	(5)
W	213	Artificial or Unknown found in <213> in SEQ ID (	5)
E	257	Invalid sequence data feature in <221> in SEQ ID	(6)
Ε	257	Invalid sequence data feature in <221> in SEQ ID	(6)
W	213	Artificial or Unknown found in <213> in SEQ ID (	7)
Ε	257	Invalid sequence data feature in <221> in SEQ ID	(7)

## Input Set:

# Output Set:

**Started:** 2008-01-04 18:09:22.369 **Finished:** 2008-01-04 18:09:27.470

**Elapsed:** 0 hr(s) 0 min(s) 5 sec(s) 101 ms

Total Warnings: 69
Total Errors: 72
No. of SeqIDs Defined: 75

Actual SeqID Count: 75

Error code		Error Description
E	257	Invalid sequence data feature in <221> in SEQ ID (7)
W	213	Artificial or Unknown found in <213> in SEQ ID (8)
E	257	Invalid sequence data feature in <221> in SEQ ID (8)
E	257	Invalid sequence data feature in <221> in SEQ ID (8)
W	213	Artificial or Unknown found in <213> in SEQ ID (9)
E	257	Invalid sequence data feature in <221> in SEQ ID (9)
E	257	Invalid sequence data feature in <221> in SEQ ID (9)
W	213	Artificial or Unknown found in <213> in SEQ ID (10)
E	257	Invalid sequence data feature in <221> in SEQ ID (10)
E	257	Invalid sequence data feature in <221> in SEQ ID (10) This error has occured more than 20 times, will not be displayed
W	213	Artificial or Unknown found in <213> in SEQ ID (11)
W	213	Artificial or Unknown found in <213> in SEQ ID (12)
W	213	Artificial or Unknown found in <213> in SEQ ID (13)
W	213	Artificial or Unknown found in <213> in SEQ ID (14)
W	213	Artificial or Unknown found in <213> in SEQ ID (15)
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W	213	Artificial or Unknown found in <213> in SEQ ID (17)
W	213	Artificial or Unknown found in <213> in SEQ ID (18)
W	213	Artificial or Unknown found in <213> in SEQ ID (19)
W	213	Artificial or Unknown found in <213> in SEQ ID (20) This error has occured more than 20 times, will not be displayed
W	402	Undefined organism found in <213> in SEQ ID (66)

### Input Set:

# Output Set:

**Started:** 2008-01-04 18:09:22.369

Finished: 2008-01-04 18:09:27.470

**Elapsed:** 0 hr(s) 0 min(s) 5 sec(s) 101 ms

Total Warnings: 69

Total Errors: 72

No. of SeqIDs Defined: 75

Actual SeqID Count: 75

Error code		Error Description				
W	402	Undefined organism found in <213> in SEQ ID (68	)			
W	402	Undefined organism found in $\langle 213 \rangle$ in SEQ ID (72	)			
W	402	Undefined organism found in <213> in SEQ ID (73	<b>(</b> )			

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      McKeating, Jane A.
       Dash, Srikanta
       Coy, David H.
<120> FLAVIVIRUS FUSION INHIBITORS
<130> 12920.0014.PCUS00
<140> 10532480
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       carbohydrate
<400> 1
Tyr Gln Val Arg Asn Ser Ser Gly Leu Tyr His Val Thr Asn Asp Cys
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Pro Asn Ser Ser Ile Val Tyr Glu Ala Ala Asp Ala Ile Leu
20 25 30

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       carbohydrate
<400> 2
Cys Ser Ala Leu Tyr Trp Val Gly Asp Leu Cys Gly Ser Val Phe Leu
                5
                                    10
                                                        15
Val Gly Gln Leu Phe Thr Phe Ser Pro Arg Arg His Trp Thr Thr Gln
            2.0
                                2.5
Asp Cys
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       t-butyloxycarbonyl, lipid, polyethylene glycol, or carbohydrate
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       macromolecular, t-butyloxycarbonyl, lipid, polyethyleneglycol, or
       carbohydrate
<400> 3
Ser Pro Arg Arg His Trp Thr Thr Gln Asp Cys Asn Cys Ser Ile Tyr
               5
                                   10
Pro Gly His Ile Thr Gly His Arg Met Ala Trp Asp Met Met Asn
                                25
Trp Ser Pro Thr
       35
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       carbohydrate
<400> 4
Met Met Met Asn Trp Ser Pro Thr Ala Ala Leu Leu Arg Ile Pro Gln
               5
                                   10
                                                        15
Ala Ile Met Asp Met Ile Ala Gly Ala His Trp Gly Val Leu Ala Gly
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25

30

20

35 40

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<210>
       5
<211> 30
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       group or one of the following groups: amido,
                                                    hydrophobic,
       macromolecular, t-butyloxycarbonyl, lipid, polyethyleneglycol, or
       carbohydrate
<400> 5
Arg Val Thr Asp Pro Asp Thr Asn Thr Thr Ile Leu Thr Asn Cys Cys
                5
                                    1.0
                                                        15
Gln Arg Asn Gln Val Ile Tyr Cys Ser Pro Ser Thr Cys Leu
            20
                                25
                                                    30
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<400> 6

Arg Asp Phe Val Glu Gly Val Ser Gly Gly Ser Trp Val Asp Ile Val 1 5 10 15

Leu Glu His Gly Ser Cys Val Thr Thr Met Ala Lys Asn Lys Pro Thr 20 25 30

Leu Asp Phe

35

<210> 7

<211> 35

<212> PRT

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<400> 7

Arg Asp Phe Ile Glu Gly Ala Ser Gly Ala Thr Trp Val Asp Leu Val 1 5 10 15

Leu Glu Gly Asp Ser Cys Leu Thr Ile Met Ala Asn Asp Lys Pro Thr
20 25 30

Leu Asp Val

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       macromolecular, t-butyloxycarbonyl, lipid, polyethyleneglycol, or
       carbohydrate
<400> 8
Arg Asp Phe Ile Glu Gly Val His Gly Gly Thr Trp Val Ser Ala Thr
                5
                                    10
                                                        15
Leu Glu Gln Asp Lys Cys Val Thr Val Met Ala Pro Asp Lys Pro Ser
                                25
                                                    30
Leu Asp Ile
        35
<210> 9
<211> 35
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       hydrophobic, macromolecular, carbobenzoxyl, dansyl,
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t-butyloxycarbonyl, lipid, polyethylene glycol, or carbohydrate

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      group or one of the following groups: amido, hydrophobic,
      macromolecular, t-butyloxycarbonyl, lipid, polyethyleneglycol, or
      carbohydrate
<400> 9
Arg Asp Phe Leu Glu Gly Val Ser Gly Ala Thr Trp Val Asp Leu Val
               5
                                    10
Leu Glu Gly Asp Ser Cys Val Thr Ile Met Ser Lys Asp Lys Pro Thr
           20
                                25
                                                    30
Ile Asp Val
        35
<210> 10
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      macromolecular, t-butyloxycarbonyl, lipid, polyethyleneglycol, or
      carbohydrate
<400> 10
Gly Gln Leu Ala Cys Lys Glu Asp Tyr Arg Tyr Ala Ile Ser Ser Thr
                5
                                    10
```

Asn Glu Ile Gly Leu Cly Ala Gly Gly Leu Thr Thr Trp Lys
20 25 30

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Glu Tyr Asn
        35
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      macromolecular, t-butyloxycarbonyl, lipid, polyethyleneglycol, or
      carbohydrate
<400> 11
Gly His Leu Asp Cys Lys Pro Glu Phe Ser Tyr Ala Ile Ala Lys Asp
               5
                                   10
                                                        15
Glu Arg Ile Gly Gln Leu Gly Ala Glu Gly Leu Thr Thr Trp Lys
           20
                                25
Glu Tyr Ser
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<223> The amino-terminal amino acid residue comprises an amino group or

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is modified to contain one of the following groups: acetyl,
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       macromolecular, t-butyloxycarbonyl, lipid, polyethyleneglycol, or
       carbohydrate
<400> 12
Gly Glu Phe Ala Cys Arg Glu Asp His Arg Tyr Ala Leu Ala Lys Thr
                5
                                    10
                                                        15
Lys Glu Ile Gly Pro Leu Gly Ala Glu Ser Leu Thr Thr Trp Thr
                                25
            20
Asp Tyr Gln
        35
<210> 13
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       macromolecular, t-butyloxycarbonyl, lipid, polyethyleneglycol, or
       carbohydrate
<400> 13
Thr Cys Asp Ala Leu Asp Ile Gly Glu Leu Cys Gly Ala Cys Val Leu
                                    10
                                                        15
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Val Gly Asp Trp Leu Val Arg His Trp Leu Ile His Ile Asp Leu Asn 20 25 30

Glu Thr

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<211> 34

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Lys Arg Phe Val Cys Lys His Ser Met Val Asp Arg Gly Trp Gly Asn 1 5 10 15

Gly Cys Gly Leu Phe Gly Lys Gly Gly Ile Val Thr Cys Ala Met Phe 20 25 30

Thr Cys

<210> 15

<211> 34

<212> PRT

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      carbohydrate
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Ser Ser Tyr Val Cys Lys Gln Gly Phe Thr Asp Arg Gly Trp Gly Asn
                5
                                   10
                                                        15
Gly Cys Gly Leu Phe Gly Lys Gly Ser Ile Asp Thr Cys Ala Lys Phe
           20
                                25
Ser Cys
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      macromolecular, t-butyloxycarbonyl, lipid, polyethyleneglycol, or
      carbohydrate
<400> 16
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1 5 10 15

Gly Cys Gly Leu Phe Gly Lys Gly Ser Ile Val Ala Cys Ala Lys Phe 20 25 30

Thr Cys

<210> 17

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Gly Cys Gly Leu Phe Gly Lys Gly Ser Ile Asp Thr Cys Ala Lys Phe 20 25 30

Ala Cys

<210> 18

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- <221> MOD\_RES
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